

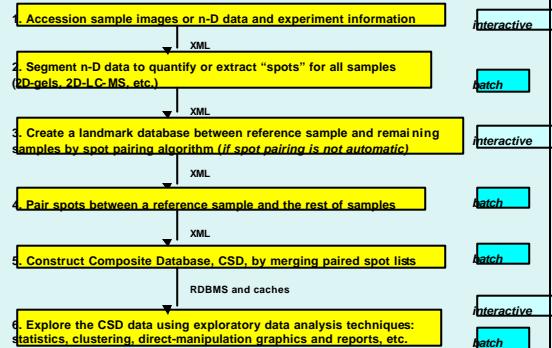
Overview: Accession

Data Accessioning Program

<http://open2dprot.sourceforge.net/Accession>

Revised: 03-01-2004, P. Lemkin

Basic Open n-D Analysis Pipeline



Introduction

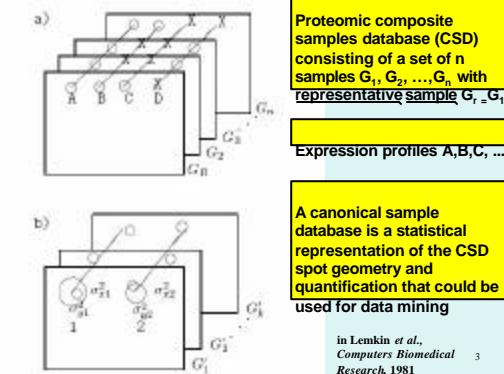
- Data must be accessioned or entered into the program database. This consists of several activities:
- 1. Entering the name of the sample and sample information
- 2. Entering the Region Of Interest of sample
- 3. Entering sample calibration info if any

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Initial Open n-D Data-Mining Tools

- Accession n-D sample images or n-D data and experiment data
- Quantify 'spots' from sample images or peptide clusters
- Pair spots between samples and a reference sample
- Construct composite sample database for exploratory data analysis
- Manage subsets of proteins in the database
- Manage replicate samples and condition sets of samples
- Analyze expression profiles for multiple conditions
- Data-filter protein sets by statistics, clustering, set membership
- Direct-manipulation of data in graphics, spreadsheets
- Integrate R language statistical, clustering, classifiers, class prediction, and other methods
- Integrate access to Internet proteomic/genomic/function data servers for user-specified protein sets

Composite Samples Database (CSD) Paradigm



Open2Dprot Pipeline Subprojects

Open2Dprot pipeline subprojects

Open2Dprot consists of a series of coordinated Open2Dprot pipeline accession modules. By using XML as the 'glue' between modules, it is possible to integrate alternate modules at the various pipeline steps. As pipeline modules and alternate modules become available, they will be added to this table. We encourage the addition of alternative pipeline processing modules which will be added to this table.

We will be using a resource, [Open2Dprot Wiki](#), in the Open2Dprot pipeline modules. This will help ensure that they are the most up-to-date, data structures and XML data interchange formats.

Subproject Name	Resource Documentation	Download (PDF)	XML documents	Version	Publication	Status	Project owner
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot
bio2dprot	bio2dprot	bio2dprot.pdf	bio2dprot.xml	bio2dprot	bio2dprot	bio2dprot	bio2dprot

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01-09-2005

Associated or Related Projects

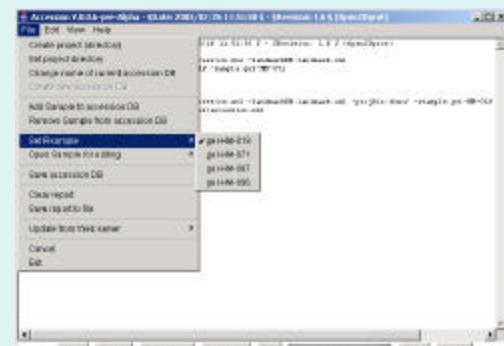
We had added some additional non-pipeline open source projects that may use similar data or function software modules. They may be useful for performing other types of analysis on data used by OptiMLDpro or alternate types of analysis.

Category	Project Name	Download	Documentation	Overview (PDF)	PDF documents	Versions	Publication History	Status
Code	Code	Code	Code	Code	Code	Code	Code	Code
DATA	bio2rdf	bio2rdf	bio2rdf	MICepr	MICepr	MICepr	MICepr	MICepr
DATA	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf
DATA	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf	bio2rdf

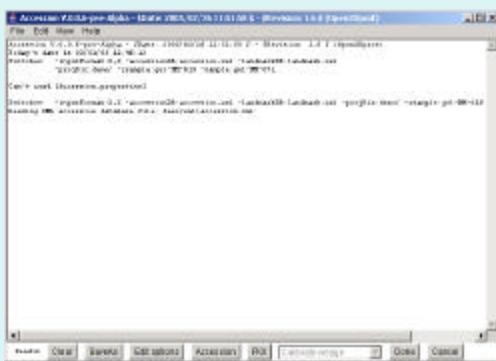
09-12-2004

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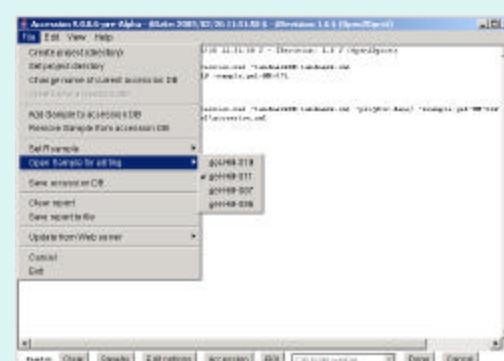
File Menu - Select Reference Sample



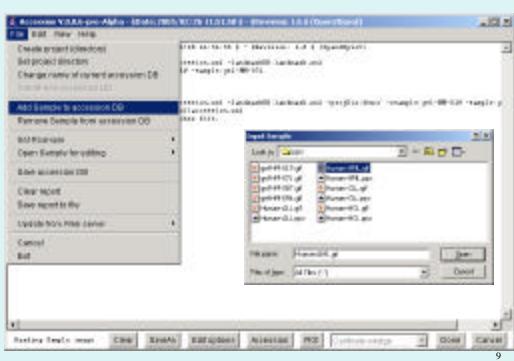
Main Window



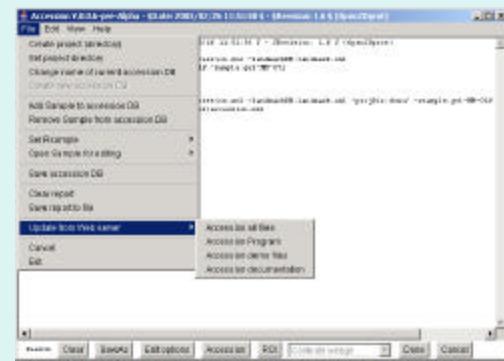
File Menu - Select Current Sample

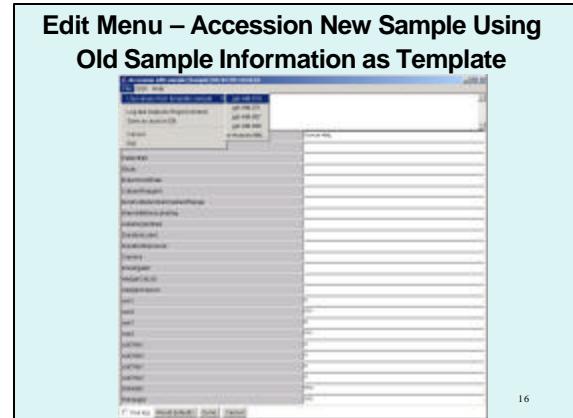
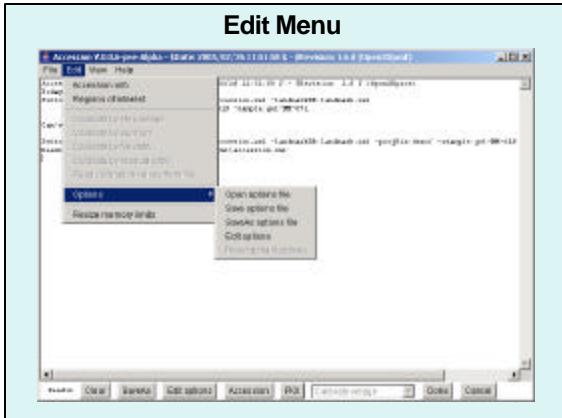


File Menu – Add New Sample

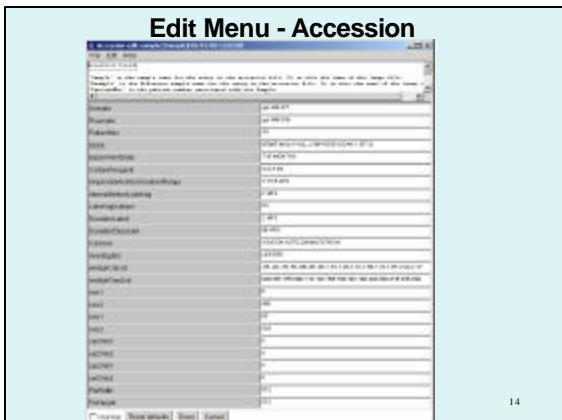


File Menu - Update From Web Server

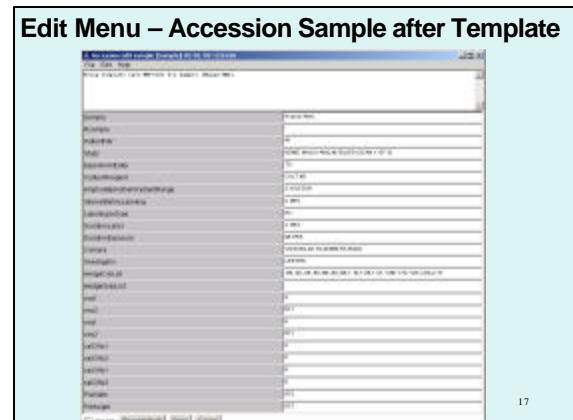




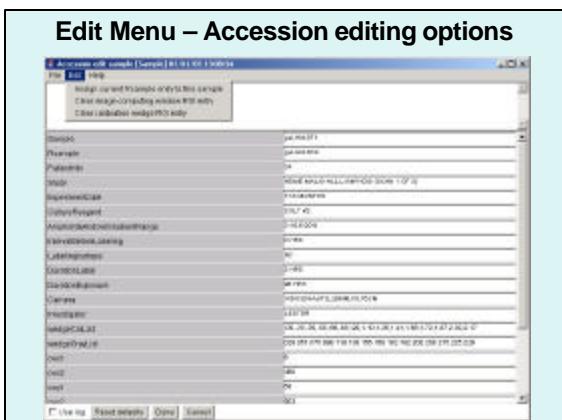
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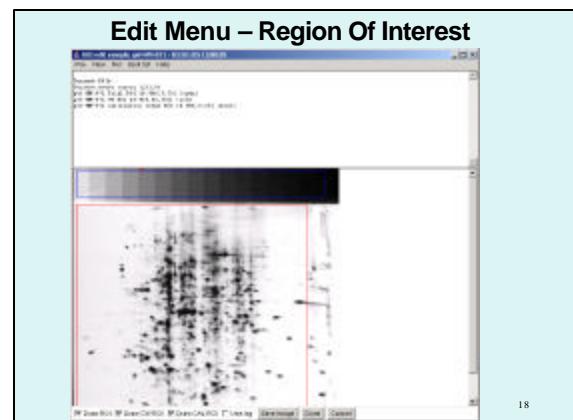
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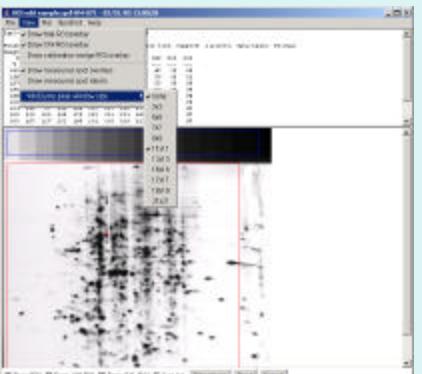


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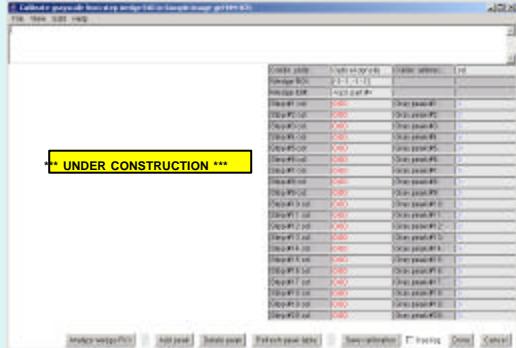
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Edit Menu – Region Of Interest overlay views



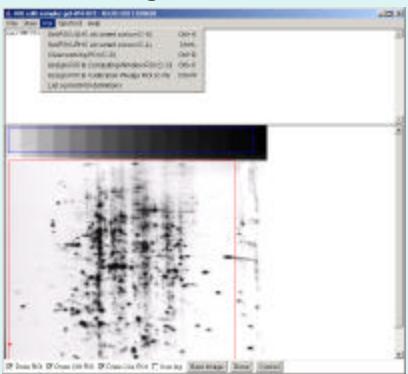
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Edit Menu – Calibrate Grayscale – OD, etc.



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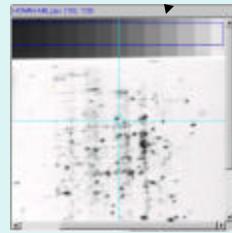
Edit Menu – Region Of Interest commands



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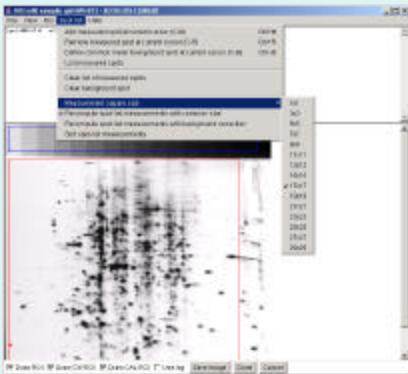
Flicker - Calibrating Grayscale with a ND Step-Wedge

1. The ND step wedge must be scanned with the image and the corresponding OD values known for each step
2. A region of interest (ROI) is overlaid on the step step-wedge
3. The ND wedge calibration wizard is invoked to analyze the data and estimate the calibration



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Edit Menu – Region Of Interest Spot List

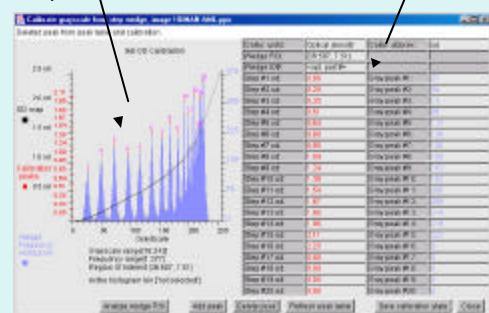


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Flicker - Calibration Curve made from ND-Wedge Data

ROI histogram_peaks found and extrapolated calibration curve

OD vs gray-peaks table



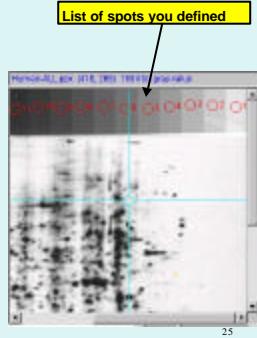
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Flicker- Calibrating Grayscale with a Spot List of Calibrated Data

1. The image must contain calibrated regions with known concentrations or corresponding OD values known for each spot

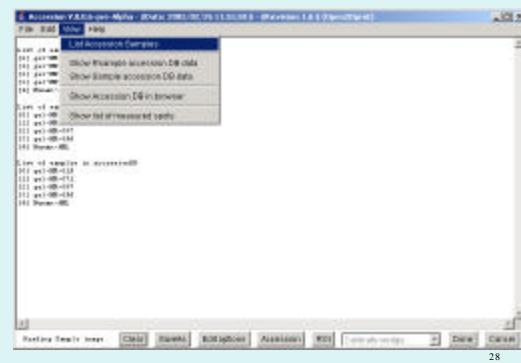
2. You define a set of spots using (C-M) or (ALT-click)

3. The Spot List Calibration wizard is invoked to analyze the data and estimate the calibration



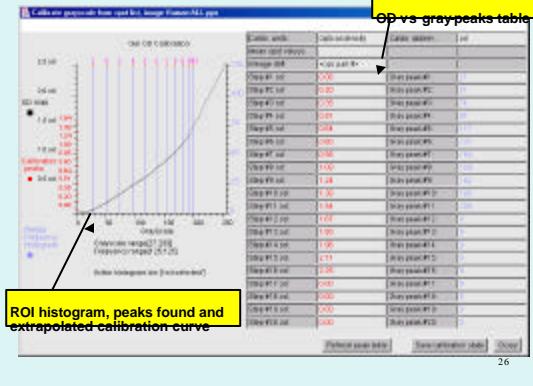
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View Menu



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Flicker - Calibrating Grayscale from Spot List Data

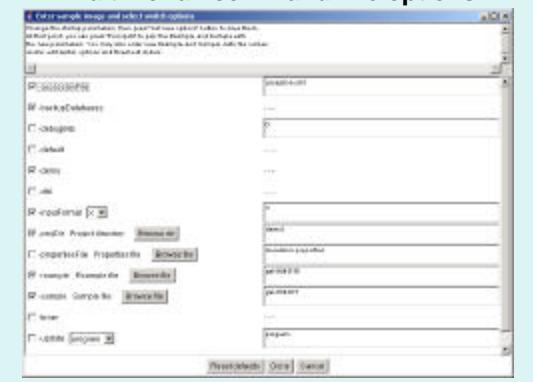


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View Menu - Show Current Sample Acc DB Data

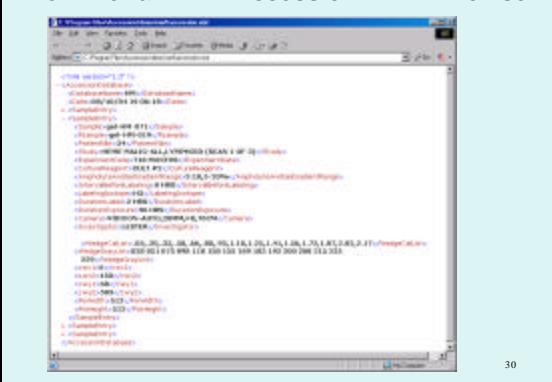


Edit Menu - command line options

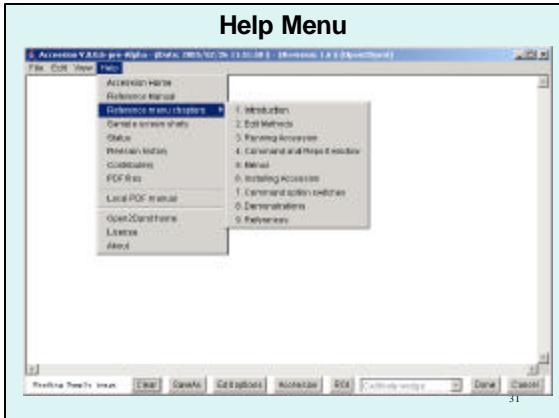


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View Menu - XML Accession DB in Browser



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Summary

- Accession is an open-source sample accessioning Java program freely available at
<http://open2dprot.sourceforge.net/Accession>
- Useful for adding sample experiment information, regions of interest and grayscale calibration (if available).
- It will be used as one of the step [1] alternative modules in the analysis pipeline in the Open2Dprot project at
<http://open2dprot.sourceforge.net>

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